STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING
   ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information
Center (STIC) detected errors when processing the following computer readable
form:

Application Serial Number: _10/557,992_

Source: _PCT/10_

Date Processed by STIC: _12/2/05_

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:
1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE
   APPLICANT, WITH A NOTICE TO COMPLY or,
2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A
   NOTICE TO COMPLY
FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT
MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER
VERSION 4.2.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND
TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:
   http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is
a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.
Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.
Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the
United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm> , EFS Submission
   User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
   U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street,
   Alexandria, VA 22314

Revised 01/24/05

PCT

```
RAW SEQUENCE LISTING              DATE: 12/02/2005
PATENT APPLICATION:  US/10/557,992   TIME: 11:33:25

         Input Set : A:\PTO.SR.txt
         Output Set: N:\CRF4\12022005\J557992.raw

   3 <110> APPLICANT: Japan Science and Technology Corporation
   5 <120> TITLE OF INVENTION: Probe for visualizing protein interaction and method of
analyzing protein-
   6        protein interaction using the same
   8 <130> FILE REFERENCE: 04F025PCT
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/557,992
C--> 10 <141> CURRENT FILING DATE: 2005-11-22
E--> 10 <160> NUMBER OF SEQ ID NOS:
```

(handwritten) insert
<1407 10/557,992
<1417 2005-11-22

(handwritten) Does Not Comply ~ see pp 1-6
Corrected Diskette Needed

(handwritten) ⑥7 (p.4)

ERRORED SEQUENCES

(handwritten right margin) Per 1.822 of Sequence Rules, number the amino acids under every — 5 amino = acids only

```
   12 <210> SEQ ID NO: 1
   13 <211> LENGTH: 300
   14 <212> TYPE: PRT
   15 <213> ORGANISM: Renilla reniformis
W--> 16 <400> SEQUENCE: 1
   17    Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile
   18    1               5                  10                  15
   19    Thr Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu
E--> 20    16              20                  25                  30
   21    Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn
E--> 22    31              35                  40                  45
   23    Ala Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp
E--> 24    46              50                  55                  60
   25    Arg His Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile
E--> 26    61              65                  70                  75
   27    Pro Asp Leu Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly
E--> 28    76              80                  85                  90
   29    Ser Tyr Arg Leu Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp Phe
E--> 30    91              95                  100                 105
   31    Glu Leu Leu Asn Leu Pro Lys Lys Ile Ile Phe Val Gly His Asp
E--> 32    106             110                 115                 120
   33    Trp Gly Ala Cys Leu Ala Phe His Tyr Cys Tyr Glu His Gln Asp
E--> 34    121             125                 130                 135
   35    Lys Ile Lys Ala Ile Val His Ala Glu Ser Val Val Asp Val Ile
E--> 36    136             140                 145                 150
   37    Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu Glu Asp Ile Ala Leu
E--> 38    151             155                 160                 165
   39    Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu Glu Asn Asn Phe
E--> 40    166             170                 175                 180
   41    Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg Lys Leu Glu
E--> 42    181             185                 190                 195
   43    Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu Lys Gly
```

(handwritten) delete these nos — (marks beside lines 18, 19)

(handwritten) see p. 2

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : A:\PTO.SR.txt
Output Set: N:\CRF4\12022005\J557992.raw

```
E--> 44    ̶1̶9̶6̶            200                 205                 210
     45   Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro Leu
E--> 46    ̶2̶1̶1̶            215                 220                 225
     47   Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr
E--> 48    ̶2̶2̶6̶            230                 235                 240
     49   Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe Ile
E--> 50    ̶2̶4̶1̶            245                 250                 255
     51   Glu Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala
E--> 52    ̶2̶5̶6̶            260                 265                 270
     53   Lys Lys Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His
E--> 54    ̶2̶7̶1̶            275                 280                 285
     55   Phe Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Asn Tyr Ile Gln
E--> 56    ̶2̶8̶6̶      .      290→               295→               300→
     59   <210> SEQ ID NO: 2
     60   <211> LENGTH: 13
     61   <212> TYPE: PRT
     62   <213> ORGANISM: Artificial Sequence
W--> 63   <220>
     63   <223> OTHER INFORMATION: Synthesized Oligopeptide
E--> 64   <400> SEQUENCE: 2
     65    Thr Glu Glu Ala Tyr Met Lys Met Asp Leu Gly Pro Gly
     66    1               5                   10
     69   <210> SEQ ID NO: 3
     70   <211> LENGTH: 300
     71   <212> TYPE: PRT
     72   <213> ORGANISM: Renilla reniformis
W--> 73   <400> SEQUENCE: 3
     74    Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile
     75    1               5                   10                  15
     76    Thr Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu
E--> 77    ̶1̶6̶            20                  25                  30
     78    Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn
E--> 79    ̶3̶1̶            35                  40                  45
     80    Ala Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp
E--> 81    ̶4̶6̶            50                  55                  60
     82    Arg His Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile
E--> 83    ̶6̶1̶            65                  70                  75
     84    Pro Asp Leu Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly
E--> 85    ̶7̶6̶            80                  85                  90
     86    Ser Tyr Arg Leu Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp Phe
E--> 87    ̶9̶1̶            95                  100                 105
     88    Glu Leu Leu Asn Leu Pro Lys Lys Ile Ile Phe Val Gly His Asp
E--> 89    ̶1̶0̶6̶           110                 115                 120
     90    Trp Gly Ala Ala Leu Ala Phe His Tyr Cys Tyr Glu His Gln Asp
E--> 91    ̶1̶2̶1̶           125                 130                 135
     92    Lys Ile Lys Ala Ile Val His Ala Glu Ser Val Val Asp Val Ile
E--> 93    ̶1̶3̶6̶           140                 145                 150
     94    Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu Glu Asp Ile Ala Leu
E--> 95    ̶1̶5̶1̶           155                 160                 165
```

[handwritten annotations:]

290→ directly under "Asp"   295→ directly under "Met"   300→ directly under "Gln"

← insert <220> wherever <221>, <222>, or <223> is shown. <220> never has a response. It is a "header" only.

same error as Sequence 1

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : A:\PTO.SR.txt
Output Set: N:\CRF4\12022005\J557992.raw

```
        96    Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu Glu Asn Asn Phe
E-->    97    166            170            175            180
        98    Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg Lys Leu Glu
E-->    99    181            185            190            195
        100   Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu Lys Gly
E-->    101   196            200            205            210
        102   Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro Leu
E-->    103   211            215            220            225
        104   Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn Tyr
E-->    105   226            230            235            240
        106   Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe Ile
E-->    107   241            245            250            255
        108   Glu Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly Ala
E-->    109   256            260            265            270
        110   Lys Lys Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu His
E-->    111   271            275            280            285
        112   Phe Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Asn Tyr Ile Gln
        113   286            290            295            300
        116   <210> SEQ ID NO: 4
        117   <211> LENGTH: 18
        118   <212> TYPE: PRT
        119   <213> ORGANISM: Artificial Sequence
W-->    120   <220>
        120   <223> OTHER INFORMATION: Synthesized Oligopeptide
        121   <400> SEQUENCE: 4
        122   Cys Leu Ser Leu Ala Ser Asn Asn Gly Asn Gly Arg Asn Gly Ala
        123   1              5              10             15
        124   Ser Leu Glu
        125   16
        128   <210> SEQ ID NO: 5
        129   <211> LENGTH: 17
        130   <212> TYPE: PRT
        131   <213> ORGANISM: Artificial Sequence
W-->    132   <220>
        132   <223> OTHER INFORMATION: Synthesized Oligopeptide
        133   <400> SEQUENCE: 5
        134   Pro Arg Gly Asn Asn Gly Gly Asn Asn Asp Val Met Ala Ile Ala
        135   1              5              10             15
        136   Ala Asn
        137   16
        140   <210> SEQ ID NO: 6
        141   <211> LENGTH: 133
        142   <212> TYPE: PRT
        143   <213> ORGANISM: Artificial Sequence
W-->    144   <220>
        144   <223> OTHER INFORMATION: Synthesized Oligopeptide
        145   <400> SEQUENCE: 6
        146   Met Thr Ser Lys Val Tyr Asp Pro Glu Gln Arg Lys Arg Met Ile
        147   1              5              10             15
```

handwritten annotations:
E--> (lines 97, 99, 101, 103, 105, 107, 109, 111 with struck-through numbers)
directly "under Met" (at line 113)
W--> 120 <220> ← insert <220>
ØK--> 121
125 16 delete
W--> 132 <220> ← insert <220>
137 16 delete
W--> 144 <220> ← insert <220>
145

RAW SEQUENCE LISTING   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:25

Input Set : A:\PTO.SR.txt
Output Set: N:\CRF4\12022005\J557992.raw

```
         148     Thr Gly Pro Gln Trp Trp Ala Arg Cys Lys Gln Met Asn Val Leu
E--> 149     16              20              25              30
         150     Asp Ser Phe Ile Asn Tyr Tyr Asp Ser Glu Lys His Ala Glu Asn
E--> 151     31              35              40              45
         152     Ala Val Ile Phe Leu His Gly Asn Ala Ala Ser Ser Tyr Leu Trp
E--> 153     46              50              55              60
         154     Arg His Val Val Pro His Ile Glu Pro Val Ala Arg Cys Ile Ile
E--> 155     61              65              70              75
         156     Pro Asp Leu Ile Gly Met Gly Lys Ser Gly Lys Ser Gly Asn Gly
E--> 157     76              80              85              90
         158     Ser Cys Leu Ser Leu Ala Ser Asn Asn Gly Asn Gly Arg Asn Gly
E--> 159     91              95              100             105
         160     Ala Ser Leu Glu Thr Glu Glu Tyr Met Lys Met Asp Leu Gly Pro
E--> 161     106             110             115             120
         162     Gly Thr Arg Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
E--> 163     121         125→ directly  "  130→ directly
```

166 <210> SEQ ID NO: 7
167 <211> LENGTH: 352
168 <212> TYPE: PRT
169 <213> ORGANISM: Artificial Sequence
W--> 170 <220> ~~insert <2207~~
170 <223> OTHER INFORMATION: Synthesized Oligopeptide
E OK 171 <400> SEQUENCE: 7

```
         172     Met Asp Ala Glu Trp Tyr Trp Gly Asp Ile Ser Arg Glu Glu Val
         173     1               5               10              15
         174     Asn Glu Lys Leu Arg Asp Thr Ala Asp Gly Thr Phe Leu Val Arg
E--> 175     16              20              25              30
         176     Asp Ala Ser Thr Lys Met His Gly Asp Tyr Thr Leu Thr Leu Arg
E--> 177     31              35              40              45
         178     Lys Gly Gly Asn Asn Lys Leu Ile Lys Ile Phe His Arg Asp Gly
E--> 179     46              50              55              60
         180     Lys Tyr Gly Phe Ser Asp Pro Leu Thr Phe Asn Ser Val Val Glu
E--> 181     61              65              70              75
         182     Leu Ile Asn His Tyr Arg Asn Glu Ser Leu Ala Gln Tyr Asn Pro
E--> 183     76              80              85              90
         184     Lys Leu Asp Val Lys Leu Leu Tyr Pro Val Ser Lys Tyr Gln Gln
E--> 185     91              95              100             105
         186     Pro Arg Gly Asn Asn Gly Gly Asn Asn Asp Val Met Ala Ile Ala
E--> 187     106             110             115             120
         188     Ala Asn Tyr Arg Leu Leu Asp His Tyr Lys Tyr Leu Thr Ala Trp
E--> 189     121             125             130             135
         190     Phe Glu Leu Leu Asn Leu Pro Lys Lys Ile Ile Phe Val Gly His
E--> 191     136             140             145             150
         192     Asp Trp Gly Ala Cys Leu Ala Phe His Tyr Ser Tyr Glu His Gln
E--> 193     151             155             160             165
         194     Asp Lys Ile Lys Ala Ile Val His Ala Glu Ser Val Val Asp Val
E--> 195     166             170             175             180
         196     Ile Glu Ser Trp Asp Glu Trp Pro Asp Ile Glu Glu Asp Ile Ala
E--> 197     181             185             190             195
```

directly under Gln

directly under "Glu"

```
        198    Leu Ile Lys Ser Glu Glu Gly Glu Lys Met Val Leu Glu Asn Asn
E--> 199    196                 200              ·          205                    210
        200    Phe Phe Val Glu Thr Met Leu Pro Ser Lys Ile Met Arg Lys Leu
E--> 201    211                 215                         220                    225
        202    Glu Pro Glu Glu Phe Ala Ala Tyr Leu Glu Pro Phe Lys Glu Lys
E--> 203    226                 230                         235                    240
        204    Gly Glu Val Arg Arg Pro Thr Leu Ser Trp Pro Arg Glu Ile Pro
E--> 205    241                 245                         250                    255
        206    Leu Val Lys Gly Gly Lys Pro Asp Val Val Gln Ile Val Arg Asn
E--> 207    256                 260                         265                    270
        208    Tyr Asn Ala Tyr Leu Arg Ala Ser Asp Asp Leu Pro Lys Met Phe
E--> 209    271                 275                         280                    285
        210    Ile Glu Ser Asp Pro Gly Phe Phe Ser Asn Ala Ile Val Glu Gly
E--> 211    286                 290                         295                    300
        212    Ala Lys Lys Phe Pro Asn Thr Glu Phe Val Lys Val Lys Gly Leu
E--> 213    301                 305                         310                    315
        214    His Phe Ser Gln Glu Asp Ala Pro Asp Glu Met Gly Lys Tyr Ile
E--> 215    316                 320                         325                    330
        216    Lys Ser Phe Val Glu Arg Val Leu Lys Asn Glu Gln Pro Arg Asp
E--> 217    331                 335                         340                    345
        218    Tyr Lys Asp Asp Val Val Lys
        219    346                 350
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 12/02/2005
PATENT APPLICATION: **US/10/557,992**   TIME: 11:33:26

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

Invalid Line Length:
The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

VERIFICATION SUMMARY   DATE: 12/02/2005
PATENT APPLICATION: US/10/557,992   TIME: 11:33:26

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

```
L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:16 M:283 W: Missing Blank Line separator, <400> field identifier
L:20 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:22 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:24 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:26 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:28 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:30 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:32 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:34 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:36 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:38 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:40 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:42 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:44 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:46 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:48 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:50 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:52 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:54 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:56 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:1
L:63 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:64 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:2
L:64 M:283 W: Missing Blank Line separator, <400> field identifier
L:73 M:283 W: Missing Blank Line separator, <400> field identifier
L:77 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:79 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:81 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:83 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:85 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:87 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:89 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:91 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:93 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:95 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:97 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:99 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:101 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:103 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:105 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:107 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:109 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:111 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:3
L:120 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:121 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:4
L:121 M:283 W: Missing Blank Line separator, <400> field identifier
L:132 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:5
```

VERIFICATION SUMMARY   DATE: 12/02/2005
PATENT APPLICATION: **US/10/557,992**   TIME: 11:33:26

Input Set : **A:\PTO.SR.txt**
Output Set: **N:\CRF4\12022005\J557992.raw**

```
L:133 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:5
L:133 M:283 W: Missing Blank Line separator, <400> field identifier
L:144 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:6
L:145 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:6
L:145 M:283 W: Missing Blank Line separator, <400> field identifier
L:149 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:151 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:153 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:155 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:157 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:159 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:161 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:163 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:6
L:170 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:171 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:7
L:171 M:283 W: Missing Blank Line separator, <400> field identifier
L:175 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:177 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:179 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:181 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:183 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:185 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:187 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:189 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:191 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:193 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:195 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:197 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:199 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:201 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:203 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:205 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:207 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:209 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:211 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:213 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:215 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:217 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:7
L:10 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (6) Counted (7)
```